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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/577,827	03/26/2007	Kazumi Koga	BY0033P	9797
MERCK AND	7590 05/01/200 CO., INC		EXAMINER	
PO BOX 2000			LANDSMAN, ROBERT S	
RAHWAY, NJ 07065-0907			ART UNIT	PAPER NUMBER
			1647	
			MAIL DATE	DELIVERY MODE
			05/01/2008	PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

	Application No.	Applicant(s)					
	10/577,827	KOGA ET AL.					
Office Action Summary	Examiner	Art Unit					
	ROBERT LANDSMAN	1647					
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address					
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 16(a). In no event, however, may a reply be tinuity 17 price of the community of	N. nely filed the mailing date of this communication. D (35 U.S.C. § 133).					
Status							
1) Responsive to communication(s) filed on							
• • • • • • • • • • • • • • • • • • • •	action is non-final.						
3) Since this application is in condition for allowan		esecution as to the merits is					
	closed in accordance with the practice under <i>Ex parte Quayle</i> , 1935 C.D. 11, 453 O.G. 213.						
Disposition of Claims							
4)⊠ Claim(s) <u>2-6,8 and 9</u> is/are pending in the appli	cation.						
, ,	4a) Of the above claim(s) is/are withdrawn from consideration.						
5)⊠ Claim(s) <u>2</u> is/are allowed.							
6)⊠ Claim(s) <u>3,8 and 9</u> is/are rejected.							
7) Claim(s) <u>4-6</u> is/are objected to.							
8) Claim(s) are subject to restriction and/or	election requirement.						
Application Papers	·						
9)⊠ The specification is objected to by the Examine	•						
10) ☐ The drawing(s) filed on <u>01 May 2006</u> is/are: a)		ov the Evaminer					
Applicant may not request that any objection to the c	_ · · · · · · ·	•					
Replacement drawing sheet(s) including the correcti	- , , , , , , , , , , , , , , , , , , ,	, ,					
11) The oath or declaration is objected to by the Ex	• • • • • • • • • • • • • • • • • • • •	, ,					
Priority under 35 U.S.C. § 119	animor. Note the attached emoc	7.00.01.01.01.11.1.10.102.					
<u> </u>							
 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some * c) None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received. 							
Attachment(s)							
1) Notice of References Cited (PTO-892) 2) Notice of Draftsperson's Patent Drawing Review (PTO-948)	4) Interview Summary Paper No(s)/Mail Da						
2)	5) Notice of Informal F						
Paper No(s)/Mail Date 10/29/07.	6)						

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DETAILED ACTION

1. Formal Matters

A. Claims 1-9 are pending in the application and are the subject of this Office Action.

2. Information Disclosure Statement

A. 37 CFR 1.98(b)(5) states -

Each publication listed in an information disclosure statement must be identified by publisher, author (if any), title, relevant pages of the publication, date, and place of publication.

Therefore, the references in the 1449 filed 10/29/07 have been considered, but will not be printed on the face of the patent.

3. Specification

When a sequence is presented in a drawing, regardless of the format or the manner of presentation of that sequence in the drawing, the sequence must still be included in the Sequence Listing and a sequence identifier ("SEQ ID NO:X") must be used either in the drawing or in the Brief Description of the Drawings. See MPEP ' 2422.02. In the instant application, a sequence identifier must be used for the sequences appearing in Figure 1.

Appropriate correction is required.

4. Claim Objections

A. Claim 5 is objected to under 37 CFR 1.75(c), as being of improper dependent form for failing to further limit the subject matter of a previous claim. Applicant is required to cancel the claim(s), or amend the claim(s) to place the claim(s) in proper dependent form, or rewrite the claim(s) in independent form. Claims 1 and 5 both ultimately recite the same nucleic acid since both claims recite "consisting of SEQ ID NO:1.".

B. Claims 4-6 are objected to since they depend from canceled claim 1.

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C. Claim 9 is objected to since it would be clearer if parts (a), (b) and (c) were further labeled as "(a)(1)," "(a)(2)," "(b)(1)," etc.

5. Claim Rejections - 35 USC § 112, first paragraph - scope of enablement

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

A. Claims 2 and 8 are rejected under 35 U.S.C. 112, first paragraph, because the specification, while then being enabling for the protein of SEQ ID NO:2 and the nucleic acid of SEQ ID NO:1, does not reasonably provide enablement for nucleic acids which hybridize to SEQ ID NO:1, or for any isolated protein consisting of SEQ ID NO:2 with a substitution, deletion, addition, or insertion and having ORL1 activity. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims.

In <u>In re Wands</u>, 8USPQ2d, 1400 (CAFC 1988) page 1404, the factors to be considered in determining whether a disclosure would require undue experimentation include (1) the quantity of experimentation necessary, (2) the amount of direction or guidance presented, (3) the presence or absence of working examples, (4) the nature of the invention, (5) the state of the prior art, (6) the relative skill of those in the art, (7) the predictability or unpredictability of the art, and (8) the breadth of the claims.

First, the breadth of the claims is excessive with regard to claiming all nucleic acids which **hybridize** under stringent conditions to SEQ ID NO:1. Nucleic acid molecules which hybridize to SEQ ID NO:1 would have one or more nucleic acid substitutions, deletions, insertions and/or additions to the polynucleotide of SEQ ID NO:1 and would **encode proteins with one or more** substitutions, deletions, insertions and/or additions to the protein of SEQ ID NO:2. Claim 8 recites similar protein limitations.

Applicants provide no guidance or working examples of nucleic acid molecules which hybridize to SEQ ID NO:1, or of proteins which with one or more substitutions, deletions, insertions and/or additions to the protein of SEQ ID NO:2 and have the recited "ORL1 activity." Furthermore, Applicants do not provide a *function* of these nucleic acid molecules, or of the proteins which they encode, other than that they have "ORL1 activity." Respectfully, "binding to an antigen" could be considered an ORL1 activity.

Applicants have provided no guidance as to what critical residues are required to maintain the functional characteristics of the protein of SEQ ID NO:2. Furthermore, it is not predictable to one of ordinary skill in the art how to make a functional protein which is less than 100% identical to that of SEQ ID NO:2.

Furthermore, the scope of claim 9 is excessive with regard to Applicants claiming a method of screening any and all ORL1 proteins. The term "ORL1" is an acronym. Applicants have only taught screening the protein of SEQ ID NO:2. The specification has not disclosed how to screen all opioid "like" receptors. No guidance or working examples of opioid "like" proteins have been disclosed in the specification, nor is it predictable what the genus of proteins is which are "like" opioid receptors other than opioid receptors, especially in absence of a specific activity (e.g. binding nociceptin).

In addition, claim 9(c) recites a method which reads on **in vivo screening**. However, there is no guidance or working examples of such an assay

In summary, the breadth of the claims is excessive with regard to Applicants claiming all nucleic acids which hybridize to SEQ ID NO:1 and with regard to proteins with one or more substitutions, deletions, insertions and/or additions to the protein of SEQ ID NO:2. The same is true for the genus of proteins to be screened in claim 9. There is also a lack of guidance and working examples of these nucleic acid molecules and proteins as well as which residues are critical for protein function. These factors, along with the lack of predictability to one of ordinary skill in the art as to how to make a functional protein other that that of SEQ ID NO:2 leads the Examiner to hold that undue experimentation is necessary to practice the invention as claimed.

6. Claim Rejections - 35 USC § 112, first paragraph – written description

A. Claims 2 and 8 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

These are genus claims. Nucleic acid molecules which **hybridize** to SEQ ID NO:1 would have one or more nucleic acid substitutions, deletions, insertions and/or additions to said polynucleotides and would encode **proteins which have one or more** amino acid substitutions, deletions, insertions and/or additions to the protein of SEQ ID NO:2.

The specification and claims do not indicate what distinguishing attributes are shared by the members of the genus. Thus the scope of the claims includes numerous structural variants, and the genus is highly variant because a significant number of structural differences between genus members is permitted. Although these types of changes are routinely done in the art, the specification and claims do not provide any guidance as to what changes should be made. Structural features that could distinguish compounds in the genus from others in the nucleic acid or protein class are missing from the disclosure. No common structural attributes identify the members of the genus. The general knowledge and level of skill in the art do not supplement the omitted description because specific, not general, guidance is what is needed. Since the disclosure fails to describe the common attributes or characteristics that identify members of the genus, and because the genus is highly variant, SEQ ID NO:1, or molecules which hybridize to this nucleic acid, along with the protein of SEQ ID NO:2, alone, are insufficient to describe the genus.

Furthermore, adequate written description is lacking with regard to Applicants claiming a method of screening any and all ORL1 proteins. The term "ORL1" is an acronym. Applicants have only taught screening the protein of SEQ ID NO:2. The specification has not disclosed how to screen all opioid "like" receptors. No description of opioid "like" proteins have been disclosed in the specification, nor does the specification describe the genus of proteins which is "like" opioid receptors other than opioid receptors, especially in absence of a specific activity (e.g. binding nociceptin).

In addition, claim 9(c) recites a method which reads on **in vivo screening**. However, no such assay is described in the specification.

One of skill in the art would reasonable conclude that the disclosure fails to provide a representative number of species to describe the genus. Thus, Applicant was not in possession of the claimed genus at the time the invention was made.

7. Claim Rejections - 35 USC § 112, second paragraph

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 1-9 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

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A. Claims 3, 5, 8 and 9 recite "ORL1." This term is an acronym and should be spelled out upon first

use.

B. Claims 3, 8 and 9 are rejected since the metes and bounds of "ORL1 activity" are unknown. In

other words, it is unclear as to what "activity" is being referred.

C. Claim 8 is confusing since it recites the transitional phrase, "consisting of," as well as the protein

having an "addition," or "insertion." It is unclear how a protein consisting of SEQ ID NO:2, which is

"closed" language, can also have an "insertion," or "deletion."

8. Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis

for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on

sale in this country, more than one year prior to the date of application for patent in the United States.

A. Claims 3, 8 and 9 are rejected under 35 U.S.C. 102(b) as being anticipated by Evans et al. (US

6,432,652). The claims recite an isolated nucleic acid which hybridizes to SEQ ID NO:1 and which

encodes a protein having ORL1 activity. The claims also recite a variant of SEQ ID NO:2 as well as

methods of screening compounds for ORL1 activity. Evans teach a nucleic acid encoding an opioid

receptor which is 93.5% identical to SEQ ID NO:1 of the instant invention (see Sequence Comparison A

below). This nucleic acid has 96% similarity with SEQ ID NO:1. Therefore, it would be expected to

hybridize to SEQ ID NO:1 even under the most stringent conditions.

The protein of Evans is 98.8% identical to SEQ ID NO:2 of the instant invention (see Sequence

Comparison B below). There is, inter alia, a substitution at position 43.

Evans also teach the screening methods recited in claim 9 (column 6, lines 24-26 and Example 2).

SEQUENCE COMPARISON A

US-08-405-271A-18

; Sequence 18, Application US/08405271A

; Patent No. 6432652

; GENERAL INFORMATION:

APPLICANT: EVANS, CHRISTOPHER J.

; APPLICANT: KEITH, DUANE E.

; TITLE OF INVENTION: OPIOID RECEPTOR GENES

```
NUMBER OF SEQUENCES: 25
    CORRESPONDENCE ADDRESS:
     ADDRESSEE: MORRISON & FOERSTER
     STREET: 2000 PENNSYLVANIA AVENUE, NW, Suite 5500
     CITY: WASHINGTON
     STATE: DC
     COUNTRY:
     ZIP: 20006-1888
    COMPUTER READABLE FORM:
     MEDIUM TYPE: Floppy disk
     COMPUTER: IBM PC compatible
     OPERATING SYSTEM: PC-DOS/MS-DOS
     SOFTWARE: PatentIn Release #1.0, Version #1.30
    CURRENT APPLICATION DATA:
     APPLICATION NUMBER: US/08/405,271A
     FILING DATE: 14-MAR-1995
     CLASSIFICATION: 435
    ATTORNEY/AGENT INFORMATION:
     NAME: MURASHIGE, KATE H.
     REGISTRATION NUMBER: 29,959
     REFERENCE/DOCKET NUMBER: 22000-20526.22
    TELECOMMUNICATION INFORMATION:
     TELEPHONE: (202) 887-1500
     TELEFAX: (202) 887-0763
     TELEX: 90-4030 MRSNFOERSWSH
  INFORMATION FOR SEQ ID NO: 18:
    SEQUENCE CHARACTERISTICS:
     LENGTH: 1805 base pairs
     TYPE: nucleic acid
     STRANDEDNESS: double
     TOPOLOGY: linear
    FEATURE:
     NAME/KEY: CDS
     LOCATION: 10..1119
US-08-405-271A-18
 Query Match
                      93.5%; Score 1041; DB 3; Length 1805;
                    96.0%; Pred. No. 5.1e-193;
 Best Local Similarity
 Matches 1068; Conservative
                           0; Mismatches
                                         45;
                                                                 0:
                                             Indels
                                                       0; Gaps
          1 ATGGAGCCTCTCTCCCCGCCCCATTCTGGGAGGTTATCTACGGCAGCCACCTTCAGGGC 60
Ov
           10 ATGGAGCCCTCTTCCCCGCGCCGTTCTGGGAGGTTATCTACGGCAGCCACCTTCAGGGC 69
Db
         61 AACCTGTCCCTCAGTCCCAACCACAGTCTGCTGCCTCCGCATCTGCTGCTCAATGCC 120
Qу
           70 AACCTGTCCTCCTGAGCCCCAACCACAGTCTGCTGCCCCGCATCTGCTGCTCAATGCC 129
Db
Qу
        121 AGTCACAGCGCCTTCCTGCCCCTCGGGCTCAAGGTCACCATCGTGGGGCTCTACCTGGCC 180
            Db
        130 AGCCACGGCGCCTTCCTGCCCCTCGGGCTCAAGGTCACCATCGTGGGGCTCTACCTGGCC 189
        181 GTGTGTCGGGGGGCTCCTGGGGAACTGCCTCGTCATGTACGTCATCCTCAGGCACACC 240
Qv
            190 GTGTGTCGGAGGGCTCCTGGGGAACTGCCTTGTCATGTACGTCATCCTCAGGCACACC 249
Db
        241 AAAATGAAGACACCACAATATTTACATCTTTAACCTGGCCCTGGCAGACACTCTGGTC 300
Qv
           Db
        250 AAAATGAAGACAGCCACCAATATTTACATCTTTAACCTGGCCCTGGCCGACACTCTGGTC 309
```

Qу	301	$\tt CTGCTGACGCTGCCCTTCCAGGGCACAGACATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTGTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGCCGTTTGGGGAATCCTCCTGGGCTTCTGGGCGTTTGGGGAATCCTCCTGGGCTTCTGGGCGTTTGGGGAATCCTCCTGGGCTTCTGGGCGTTTGGGGAATCCTCCTGGGCTGTGGGAATCCTGCTGGGCTGTGGGAATCCTGGGCGGTTTGGGGAATCGTGGGGAATCCTGGGGGAATCCTGGGGCTGGGGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGGAATGCTGGGGGAATGCTGGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGGAATGCTGGGAATGCTGGAATGGAATGCTGGAATGGAATGCTGGAATGGAATGCTGGAATGGAATGCTGGAATGCTGGAATGGAATGCTGGAATGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGGAATGCTGAATGCTGAATGCTGAATGCTGAATGCTGAATGGAATGCTGAATGCTGAATGGAATGCTGAATGAA$		
Db	310	CTGCTGACGCTGCCCTTCCAGGGCACGGACATCCTCCTGGGCTTCTGGCCGTTTGGGAAT	369	
Qу	361	${\tt GCCCTGTGCAAGACAGTCATTGCCATTGACTACTACAACATGTTCACCAGCACCTTCACC}$	420	
Db	370		429	
QУ	421	$\tt CTGACTGCCATGAGTGTGGATCGCTACGTAGCCATCTGCCACCCCATCCGCGCCCTCGAC$	480	
Db	430	CTAACTGCCATGAGTGTGGATCGCTATGTAGCCATCTGCCACCCCATCCGTGCCCTCGAC	489	
Qу	481	$\tt GTCCGCACATCCAGCAAAGCCCAGGCTGTCAATGTGGCCATCTGGGCCCTGGCCTCTGTT$	540	
Db	490	GTCCGCACGTCCAGCAAAGCCCAGGCTGTCAATGTGGCCATCTGGGCCCTGGCCTCTGTT	549	
Qу	541		600	
Db	550	GTCGGTGTTCCCGTTGCCATCATGGGCTCGGCACAGGTCGAGGATGAAGAGATCGAGTGC	609	
Qу	601	$\tt CTGGTGGAGATCCCTGCCCCACAGGACTACTGGGGCCCTGTGTTTGCCGTCTGCATCTTC$	660	
Db	610	CTGGTGGAGATCCCTACCCCTCAGGATTACTGGGGCCCGGTGTTTGCCATCTTC	669	
Qу	661		720	
Db	670	CTCTTCTCCTTCATCGTCCCCGTGCTCGTCATCTCTGTCTG	729	
Qу	721	$\tt AGGCTCCGCGGAGTCCGCCTGCTCTCGGGCTCCCGGGAGAAGGACCGGAACCTGCGGCGC$	780	
Db	730	CGGCTCCGTGGAGTCCGCCTGCTCTCGGGCTCCCGAGAGAAGGACCGGAACCTGCGGCGC	789	
QУ	781	$\tt ATCACTCGGCTGGTGGTGGTGGTGGTGGTTGTTCGTGGGCTGCT$	840	
Db			849	
Qy Dl-		GTCTTTGTGCTGGTCCAAGGGCTGGGAGTGCAGCCAGGCAGCGAGACTGCCGTGGCCATT		
Db		GTCTTCGTGCTGGCCCAAGGGCTGGGGGTTCAGCCGAGCAGCGAGACTGCCGTGGCCATT	909	
QУ		CTGCGTTTCTGCACGGCCCTGGGCTACGTCAACAGCTGCCTCAACCCCATCCTCTATGCC		
Db		CTGCGCTTCTGCACGGCCCTGGGCTACGTCAACAGCTGCCTCAACCCCATCCTCTACGCC		
Qy 		TTCCTGGATGAGAACTTCAAGGCCTGCTTCCGCAAGTTCTGCTGTGCCCTGCGC		
Db		TTCCTGGATGAGAACTTCAAGGCCTGCTTCCGCAAGTTCTGCTGTGCATCTGCCCTGCGC		
~1		CGGGAGGTGCAGGTGTCCGACCGTGTGCGCAGCATTGCCAAAGATGTGGCCCTGGCCTGC		
		CGGGACGTGCAGGTGTCTGACCGCGTGCGCAGCATTGCCAAGGACGTGGCCCTGCCTG	1089	
		AAGACCTCTGAGACGGTACCGCGGCCCGCGTGA 1113		
Db	1090	AAGACCTCTGAGACGGTACCGCGGCCCGCATGA 1122		

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SEQUENCE COMPARISON B

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```
ΤD
    ABG33031 standard; protein; 370 AA.
XX
AC
    ABG33031;
XX
DT
    15-JUN-2007
               (revised)
    19-NOV-2002
DT
              (first entry)
XX
DF.
    Human opioid receptor ORL-1.
XX
OS
    Homo sapiens.
XX
PN
    US6432652-B1.
XX
    13-AUG-2002.
PD
XX
    14-MAR-1995;
                95US-00405271.
PF
XX
    (REGC ) UNIV CALIFORNIA.
PΑ
XX
PΙ
    Evans CJ, Keith DE, Edwards RH, Kaufman D;
XX
    WPI; 2002-681194/73.
DR
    N-PSDB; ABS53446.
DR
DR
    PC:NCBI; gi4505513.
    PC:SWISSPROT; P41146.
DR
DR
    PC:BIND; 177237.
CC
CC
    Revised record issued on 15-JUN-2007: Enhanced with precomputed
CC
    information from BOND.
XX
SQ
    Sequence 370 AA;
 Query Match
                      98.8%; Score 1884; DB 5; Length 370;
 Best Local Similarity
                      98.1%; Pred. No. 8.1e-200;
 Matches 363; Conservative
                            3; Mismatches
                                           4;
                                              Indels
                                                       0;
                                                          Gaps
                                                                 0:
Qу
          1 MEPLFPAPFWEVIYGSHLQGNLSLLSPNHSLLPPHLLLNASHSAFLPLGLKVTIVGLYLA 60
            1 MEPLFPAPFWEVIYGSHLQGNLSLLSPNHSLLPPHLLLNASHGAFLPLGLKVTIVGLYLA 60
Db
         61 VCVGGLLGNCLVMYVILRHTKMKTATNIYIFNLALADTLVLLTLPFQGTDILLGFWPFGN 120
QУ
            61 VCVGGLLGNCLVMYVILRHTKMKTATNIYIFNLALADTLVLLTLPFQGTDILLGFWPFGN 120
Db
        121 ALCKTVIAIDYYNMFTSTFTLTAMSVDRYVAICHPIRALDVRTSSKAQAVNVAIWALASV 180
Qу
            121 ALCKTVIAIDYYNMFTSTFTLTAMSVDRYVAICHPIRALDVRTSSKAQAVNVAIWALASV 180
Db
        181 VGVPVAIMGSAQVEDEEIECLVEIPAPQDYWGPVFAVCIFLFSFIVPVLIISVCYSLMIR 240
QУ
            181 VGVPVAIMGSAQVEDEEIECLVEIPTPQDYWGPVFAICIFLFSFIVPVLVISVCYSLMIR 240
Db
Qу
        241 RLRGVRLLSGSREKDRNLRRITRLVLVVVAVFVGCWTPVQVFVLVQGLGVQPGSETAVAI 300
            Db
        241 RLRGVRLLSGSREKDRNLRRITRLVLVVVAVFVGCWTPVQVFVLAQGLGVQPSSETAVAI 300
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9. Conclusion

A. Claim 2 is allowable. Claims 4-6 are objected to as depending from a canceled claim.

Advisory information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Robert Landsman, Ph.D. whose telephone number is (571) 272-0888. The examiner can normally be reached on M-F 10 AM -6:30 PM (eastern).

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Manjunath Rao can be reached on 571-272-0939. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

/Robert Landsman/ Primary Examiner, Art Unit 1647